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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/981,606

DATE: 05/16/2002

TIME: 10:07:34

Input Set : A:\Nano4col.app
 Output Set: N:\CRF3\05162002\I981606.raw

3 <110> APPLICANT: Rothenberg et al.
 5 <120> TITLE OF INVENTION: Mutations associated with iron disorders
 7 <130> FILE REFERENCE: 24065-004CON
 9 <140> CURRENT APPLICATION NUMBER: 09/981,606
 C--> 10 <141> CURRENT FILING DATE: 2002-10-16
 12 <150> PRIOR APPLICATION NUMBER: 09/277,457
 13 <151> PRIOR FILING DATE: 1999-03-26
 15 <160> NUMBER OF SEQ ID NOS: 30
 17 <170> SOFTWARE: PatentIn Ver. 2.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 2506
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Homo sapiens
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 28 tatgatcatg agagtcggcc tggagccat cgaactccat gggtttccag tagaatttca 240
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54 gtttttcta attggcatga aggtgtcata cagatttgca aagtttaatg gtgccttcat 1800
 55 ttgggatgct actctagttat tccagacctg aagaatcaca ataattttct acctggcttc 1860
 56 tccttgttct gataatgaaa attatgataa ggatgataaa agcacttact tcgtgtccga 1920
 57 ctcttcttag cacctactta catgcattac tgcattgact tcttacaata attctatgag 1980
 58 ataggtacta ttatccccat ttcttttta aatgaagaaa gtgaagtagg ccgggcacgg 2040
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 60 gtcaaaagag tcttaatata tataccaga tggcatgtgt ttactttatg ttactacatg 2160
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 72 <213> ORGANISM: Homo sapiens
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 78 20 25 30
 79 Leu Phe Met Gly Ala Ser Glu Gln Asp Leu Gly Leu Ser Leu Phe Glu
 80 35 40 45
 81 Ala Leu Gly Tyr Val Asp Asp Gln Leu Phe Val Phe Tyr Asp His Glu
 82 50 55 60
 83 Ser Arg Arg Val Glu Pro Arg Thr Pro Trp Val Ser Ser Arg Ile Ser
 84 65 70 75 80
 85 Ser Gln Met Trp Leu Gln Leu Ser Gln Ser Leu Lys Gly Trp Asp His
 86 85 90 95
 87 Met Phe Thr Val Asp Phe Trp Thr Ile Met Glu Asn His Asn His Ser
 88 100 105 110
 89 Lys Glu Ser His Thr Leu Gln Val Ile Leu Gly Cys Glu Met Gln Glu
 90 115 120 125
 91 Asp Asn Ser Thr Glu Gly Tyr Trp Lys Tyr Gly Tyr Asp Gly Gln Asp
 92 130 135 140
 93 His Leu Glu Phe Cys Pro Asp Thr Leu Asp Trp Arg Ala Ala Glu Pro
 94 145 150 155 160
 95 Arg Ala Trp Pro Thr Lys Leu Glu Trp Glu Arg His Lys Ile Arg Ala
 96 165 170 175
 97 Arg Gln Asn Arg Ala Tyr Leu Glu Arg Asp Cys Pro Ala Gln Leu Gln
 98 180 185 190
 99 Gln Leu Leu Glu Leu Gly Arg Gly Val Leu Asp Gln Gln Val Pro Pro
 100 195 200 205
 101 Leu Val Lys Val Thr His His Val Thr Ser Ser Val Thr Thr Leu Arg
 102 210 215 220
 103 Cys Arg Ala Leu Asn Tyr Tyr Pro Gln Asn Ile Thr Met Lys Trp Leu
 104 225 230 235 240
 105 Lys Asp Lys Gln Pro Met Asp Ala Lys Glu Phe Glu Pro Lys Asp Val

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121	245	250	255	
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124	260	265	270	
126	Pro Pro Gly Glu Glu Gln Arg Tyr Thr Cys Gln Val Glu His Pro Gly			
127	275	280	285	
129	Leu Asp Gln Pro Leu Ile Val Ile Trp Glu Pro Ser Pro Ser Gly Thr			
130	290	295	300	
132	Leu Val Ile Gly Val Ile Ser Gly Ile Ala Val Phe Val Val Ile Leu			
133	305	310	315	320
135	Phe Ile Gly Ile Leu Phe Ile Ile Leu Arg Lys Arg Gln Gly Ser Arg			
136	325	330	335	
138	Gly Ala Met Gly His Tyr Val Leu Ala Glu Arg Glu			
139	340	345		
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143	<211> LENGTH: 23			
144	<212> TYPE: DNA			
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147	<220> FEATURE:			
148	<223> OTHER INFORMATION: Description of Artificial Sequence:oligonucleotide			
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164	<400> SEQUENCE: 4			
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168	<210> SEQ ID NO: 5			
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175	primer			
177	<400> SEQUENCE: 5			
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184	<213> ORGANISM: Artificial Sequence			
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188	primer			
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Input Set : A:\Nano4col.app
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194 <210> SEQ ID NO: 7
195 <211> LENGTH: 21
196 <212> TYPE: DNA
197 <213> ORGANISM: Artificial Sequence
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208 <211> LENGTH: 22
209 <212> TYPE: DNA
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221 <211> LENGTH: 30
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229 <400> SEQUENCE: 9
230 aaaggatcca ccatggggccg gcgagccagg 30
233 <210> SEQ ID NO: 10
234 <211> LENGTH: 20
235 <212> TYPE: DNA
236 <213> ORGANISM: Artificial Sequence
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243 gtgagtctgc aggctgcgtg 20
246 <210> SEQ ID NO: 11
247 <211> LENGTH: 21
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
251 <220> FEATURE:
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255 <400> SEQUENCE: 11
256 gttccagtct tcctggcaag g 21
259 <210> SEQ ID NO: 12
260 <211> LENGTH: 22
261 <212> TYPE: DNA
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Input Set : A:\Nano4col.app
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266 primer
268 <400> SEQUENCE: 12
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273 <211> LENGTH: 21
274 <212> TYPE: DNA
275 <213> ORGANISM: Artificial Sequence
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279 primer
281 <400> SEQUENCE: 13 21
282 gttccagtct tcctggcaag g
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286 <211> LENGTH: 22
287 <212> TYPE: DNA
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300 <212> TYPE: DNA
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308 gtgtggagcc tcaacatcct g
311 <210> SEQ ID NO: 16
312 <211> LENGTH: 21
313 <212> TYPE: DNA
314 <213> ORGANISM: Artificial Sequence
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317 <223> OTHER INFORMATION: Description of Artificial Sequence:oligonucleotide
318 primer
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321 acaagacctc agacttccag c 21
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325 <211> LENGTH: 22
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327 <213> ORGANISM: Artificial Sequence
329 <220> FEATURE:
330 <223> OTHER INFORMATION: Description of Artificial Sequence:oligonucleotide
331 primer
333 <400> SEQUENCE: 17

VERIFICATION SUMMARY
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